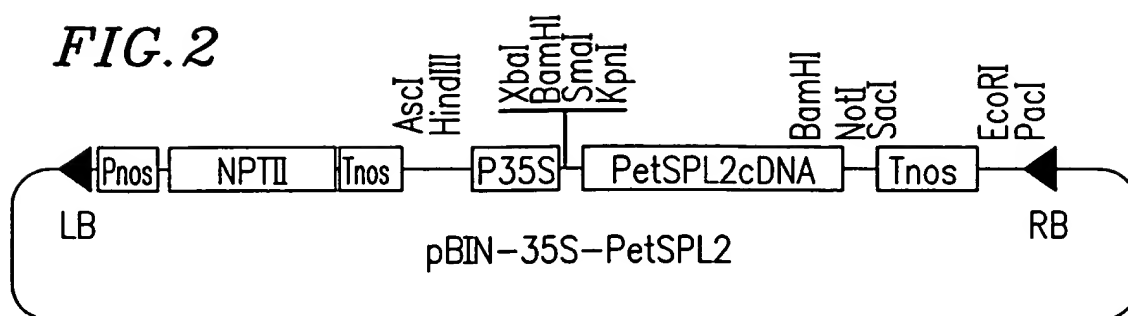


FIG. 1

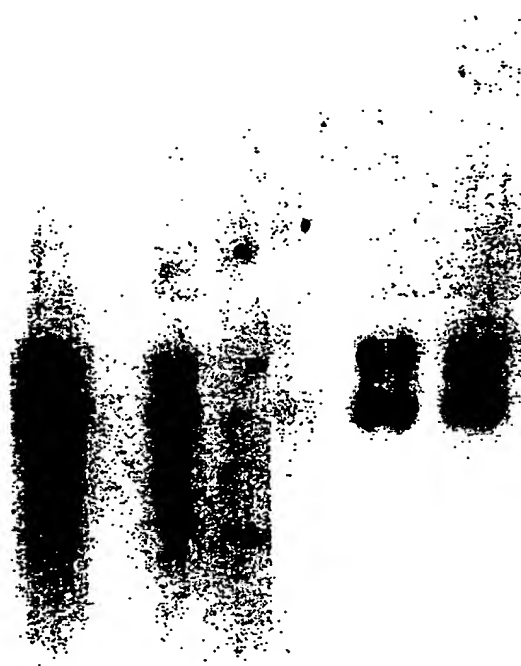
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 TGCAGTAACA CTCCTATTTA ACCCTCACAA AAAAATTACC AGAGGGCAGC AAAAAATGCT  
 TGAACATAAT TATTATACTT ACTATTAAGC TAGATTTCCCT CTTGATCTTG CTAGGTTTGA  
 CTGGAGAAAA TGGCAGGCAT GGATAGAAAC AGTTTCAACA GTAAGTACTT CAAAAACAAA  
 M A G M D R N S F N S K Y F K N K  
 AGCATCATGG CAAGACAGAT GGAGTACTTG AATAACAACA ATGGCGACAA TAACAACAAC  
 S I M A R Q M E Y L N N N N G D N N N N  
 AATAATGTTA CAAGCTCATT ACGAGATAAT TATGGAAATG AAGATCATT ACTTGGTGGG  
 N N V T S S L R D N Y G N E D H L L G G  
 CTATTCTCTT GGCCTCCAAG ATCTTATACA TGTAGCTTTT GTAAAAGGGA ATTTAGATCT  
 L F S W P P R S Y T C S F C K R E F R S  
 GCTCAAGCTC TTGGTGGACA CATGAATGTT CATAGAAGAG ATAGAGCCAT TTTGAGACAA  
 A Q A L G G H M N V H R R D R A I L R Q  
 TCACCACCTA GAGATATTAA TAGGTATTCT CTTCTAAACC TTAATCTTGA ACCAAACCCT  
 S P P R D I N R Y S L L N L N L E P N P  
 AACTTTTACC CTAGTCATAA CCCTAGTTTT TCAAGAAAAAT TCCCACCTTT TGAAATGAGG  
 N F Y P S H N P S F S R K F P P F E M R  
 AAATTAGGAA AAGGAGTTGT TCCAAACAAT CACTTGAAAA GTGCCAGAGG GCGTTTTGGA  
 K L G K G V V P N N H L K S A R G R F G  
 GTTGAGAAAA TTGACTCTTT CATGCAAGAA AAAGAATGTA CTACTACAGT GATCAAGAAG  
 V E K I D S F M Q E K E C T T T V I K K  
 TCCGAGTTTC TAAGATTGGA CTTGGGAATT GGGTTGATCA GTGAATCAAA GGAAGATTTA  
 S E F L R L D L G I G L I S E S K E D L  
 GATCTTGAAC TTCGACTGGG ATCCACTTAA CTATATCTAA TTTTACGGC ATTAAGGTTT  
 D L E L R L G S T  
 GTAAATTGAG TCGACAGCTT AGTCAAACT ACTTATGCAC TTTAATATGG CTTCTTGTGC  
 TATATTTATT TATTTTACAT GGCTGTATCT AGGTTTGCAT TTTAAGATTT AGTACCTTGT  
 CAGATTAAAA GAAAACGAAA GTTAAATTAA AAAAAA

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*FIG. 2*

*FIG. 3*

1 2 3 4 5 6 7 8 9 10 11 12 13



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*FIG. 4*



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018760-08595T60

*FIG. 5*



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868T60 08595T60